



ENTERED

PCT10

RAW SEQUENCE LISTING

DATE: 03/01/2002

PATENT APPLICATION: US/10/049,868

TIME: 15:02:30

Input Set : A:\PCT EPO 07874 Sequence listing.txt

Output Set: N:\CRF3\03012002\J049868.raw

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3 <110> APPLICANT: K. U. Leuven Research & Development
4     Deckmyn, Hans
5     Cauwenberghs, Nancy
7 <120> TITLE OF INVENTION: Cell lines, ligands and antibody fragments for use in
8     pharmaceutical compositions for preventing and treating
9     haemostasis disorders
11 <130> FILE REFERENCE: 6B4FAB
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/049,868
C--> 13 <141> CURRENT FILING DATE: 2002-02-11
13 <150> PRIOR APPLICATION NUMBER: EP00102032.0
14 <151> PRIOR FILING DATE: 2000-02-02
16 <150> PRIOR APPLICATION NUMBER: GB9918788.2
17 <151> PRIOR FILING DATE: 1999-12-10
19 <160> NUMBER OF SEQ ID NOS: 4
21 <170> SOFTWARE: PatentIn version 3.1
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25 <212> TYPE: DNA
26 <213> ORGANISM: Mus sp.
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29 <221> NAME/KEY: V_region
30 <222> LOCATION: (1)..(359)
31 <223> OTHER INFORMATION:
33 <220> FEATURE:
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35 <222> LOCATION: (1)..(24)
36 <223> OTHER INFORMATION: Vkbac primer
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39 <221> NAME/KEY: primer_bind
40 <222> LOCATION: (328)..(359)
41 <223> OTHER INFORMATION: Vkb2for primer
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44 <221> NAME/KEY: misc_feature
45 <222> LOCATION: (70)..(100)
46 <223> OTHER INFORMATION: complementary determining region one
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50 <222> LOCATION: (144)..(165)
51 <223> OTHER INFORMATION: complementary determining region two
53 <220> FEATURE:
54 <221> NAME/KEY: misc_feature
55 <222> LOCATION: (262)..(289)
56 <223> OTHER INFORMATION: complementary determining region three

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61 atgacctgca gtgccagttc aagtgttaat tacatgcact ggttccagca ggagtcgggc      120
63 accttcccca aaagaaggat ttatgacaca tccaaactgg cttctggagt ccctgctcgc      180
65 ctcaagtggca gtgggtcttg gacagaattc accctggaaa tcagtagagt gaaggctgag      240
67 gatgtgggtg tgtattactg tcaacaactt gtagagtatc cgctcacgtt cgggtgctgg      300
69 accaagctgg agctgaaacg ggctgatgct gcaccaactg tatccatctt caagcttcc      359
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73 <211> LENGTH: 363
74 <212> TYPE: DNA
75 <213> ORGANISM: Mus sp.
W--> 76 <220> FEATURE:
77 <221> NAME/KEY: V_region
78 <222> LOCATION: (1)..(363)
79 <223> OTHER INFORMATION:
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82 <221> NAME/KEY: primer_bind
83 <222> LOCATION: (1)..(24)
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87 <221> NAME/KEY: primer_bind
88 <222> LOCATION: (330)..(360)
89 <223> OTHER INFORMATION: Vhfor primer
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92 <221> NAME/KEY: misc_feature
93 <222> LOCATION: (90)..(105)
94 <223> OTHER INFORMATION: complementary determining region one
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97 <221> NAME/KEY: misc_feature
98 <222> LOCATION: (147)..(195)
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102 <221> NAME/KEY: misc_feature
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104 <223> OTHER INFORMATION: complementary determining region three
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109 acttgcaactg tctctgggat ttcattaaac agatatggtg tacactgggt tcgccagcct      120
111 ccaggaaagg gtctggagtg gctgggagta atatggactg gtggaagcac aaattataat      180
113 tcggtctctca tgtccagact gagcatcagc aaagacaact ccaagagcca agttttctta      240
115 aaaatgaaca gtctgcagac tgatgacaca gccatgtact actgtgccag agatcgatct      300
117 actatgatta cggcctatgc tatggactac tggggccaag ggaccacggt caccgtctcc      360
119 tca
122 <210> SEQ ID NO: 3
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124 <212> TYPE: PRT
125 <213> ORGANISM: Mus sp.
W--> 126 <400> SEQUENCE: 3
128 Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly

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129 1          5          10          15
132 Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser Ser Val Asn Tyr Met
133          20          25          30
136 His Trp Phe Gln Gln Glu Ser Gly Thr Phe Pro Lys Arg Arg Ile Tyr
137          35          40          45
140 Asp Thr Ser Lys Leu Ala Ser Gly Val Pro Ala Arg Leu Ser Gly Ser
141          50          55          60
144 Gly Ser Gly Thr Glu Phe Thr Leu Glu Ile Ser Arg Val Lys Ala Glu
145 65          70          75          80
148 Asp Val Gly Val Tyr Tyr Cys Gln Gln Leu Val Glu Tyr Pro Leu Thr
149          85          90          95
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156 Thr Val Ser Ile Phe Lys Leu
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162 <212> TYPE: PRT
163 <213> ORGANISM: Mus sp.
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171          20          25          30
174 Gly Val His Trp Val Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Leu
175          35          40          45
178 Gly Val Ile Trp Thr Gly Gly Ser Thr Asn Tyr Asn Ser Ala Leu Met
179          50          55          60
182 Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu
183 65          70          75          80
186 Lys Met Asn Ser Leu Gln Thr Asp Asp Thr Ala Met Tyr Tyr Cys Ala
187          85          90          95
190 Arg Asp Arg Ser Thr Met Ile Thr Ala Tyr Ala Met Asp Tyr Trp Gly
191          100          105          110
194 Gln Gly Thr Thr Val Thr Val Ser Ser
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VERIFICATION SUMMARY

DATE: 03/01/2002

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Input Set : A:\PCT EPO 07874 Sequence listing.txt

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:76 M:283 W: Missing Blank Line separator, <220> field identifier
L:126 M:283 W: Missing Blank Line separator, <400> field identifier
L:164 M:283 W: Missing Blank Line separator, <400> field identifier